FIG. 1

(SEQ ID NO: 1) gcgccgcgtc ccgcaggccg tgatgccgcc cgcgcggagg tggcccggac cgcagtgccc 60 caaqagaget ctaatggtac caagtgacag gttggettta ctgtgactcg gggacgccag 120 agetectgag aag atg tea gea ata eag gee gee tgg eea tee ggt aca 169 Met Ser Ala Ile Gln Ala Ala Trp Pro Ser Gly Thr 217 gaa tgt att gcc aag tac aac ttc cac ggc act gcc gag cag gac ctg Glu Cys Ile Ala Lys Tyr Asn Phe His Gly Thr Ala Glu Gln Asp Leu ccc ttc tgc aaa gga gac gtg ctc acc att gtg gcc gtc acc aag gac 265 Pro Phe Cys Lys Gly Asp Val Leu Thr Ile Val Ala Val Thr Lys Asp 313 ccc aac tgg tac aaa gcc aaa aac aag gtg ggc cgt gag ggc atc atc Pro Asn Trp Tyr Lys Ala Lys Asn Lys Val Gly Arg Glu Gly Ile Ile 50 cca gcc aac tac gtc cag aag cgg gag ggc gtg aag gcg ggt acc aaa Pro Ala Asn Tyr Val Gln Lys Arg Glu Gly Val Lys Ala Gly Thr Lys 361 65 409 ctc agc ctc atg cct tgg ttc cac ggc aag atc aca cgg gag cag gct Leu Ser Leu Met Pro Trp Phe His Gly Lys Ile Thr Arg Glu Gln Ala gag cgg ctt ctg tac ccg ccg gag aca ggc ctg ttc ctg gtg cgg gag 457 Glu Arg Leu Leu Tyr Pro Pro Glu Thr Gly Leu Phe Leu Val Arg Glu 95 505 age ace aac tac eee gga gae tac acg etg tge gtg age tge gae gge Ser Thr Asn Tyr Pro Gly Asp Tyr Thr Leu Cys Val Ser Cys Asp Gly aag gtg gag cac tac cgc atc atg tac cat gcc agc aag ctc agc atc 553 Lys Val Glu His Tyr Arg Ile Met Tyr His Ala Ser Lys Leu Ser Ile 135 125 130 gac gag gag gtg tac ttt gag aac ctc atg cag ctg gtg gag cac tac 601 Asp Glu Glu Val Tyr Phe Glu Asn Leu Met Gln Leu Val Glu His Tyr 649 acc tca gac gca gat gga ctc tgt acg cgc ctc att aaa cca aag gtc Thr Ser Asp Ala Asp Gly Leu Cys Thr Arg Leu Ile Lys Pro Lys Val 165 697 atg gag ggc aca gtg gcg gcc cag gat gag .ttc tac cgc agc ggc tgg Met Glu Gly Thr Val Ala Ala Gln Asp Glu Phe Tyr Arg Ser Gly Trp 180 745 gcc ctg aac atg aag gag ctg aag ctg ctg cag acc atc ggg aag ggg Ala Leu Asn Met Lys Glu Leu Lys Leu Leu Gln Thr Ile Gly Lys Gly

200

195

FIG. 1 cont.

								gtc Val		793
								gct Ala 235		841
								ctc Leu		889
								gag Glu		937
								cgg Arg		985
								tgc Cys		1033
gcc			gag			gtg		ctg Leu 315	gct	1081
								agc Ser		1129
								aag Lys		1177
								ttc Phe		1225
								atc Ile		1273
								gtc Val 395		1321
								tgc Cys		1369
								gcc Ala		1417

FIG. 1 cont.

atg cgg ccc tcc ttc cta cag ctc cga gag cag ctt gag cac atc ada Met Arg Pro Ser Phe Leu Gln Leu Arg Glu Gln Leu Glu His Ile Lys 430 435 440	1403
ace cac gag ctg cac ctg tgacggctgg cctccgcctg ggtcatgggc Thr His Glu Leu His Leu 445	1513
ctgtggggac tgaacctgga agatcatgga cetggtgcec ctgetcactg ggcccgagcc	1573
tgaactgage cecageggge tggegggeet tttteetgeg teecageetg caceeeteeg	1633
gccccgtctc tettggaccc acetgtgggg cetggggage ceaetgaggg gccagggagg	1693
aaggaggcca cggagcggga ggcagcgccc caccacgtcg ggcttccctg gcctcccgcc	1753
actcgccttc ttagagtttt attcctttcc ttttttgaga tttttttcc gtgtgtttat	1813
tttttattat ttttcaagat aaggagaaag aaagtaccca gcaaatgggc attttacaag	1873
aagtacgaat cttatttttc ctgtcctgcc cgtgagggtg ggggggaccg ggcccctctc	1933
tagggacccc tegececage eteatteece attetgtgte ceatgteecg tgteteeteg	1993
gtcgccccgt gtttgcgctt gaccatgttg cactgtttgc atgcgcccga ggcagacgtc	2053
tgtcaggggc ttggatttcg tgtgccgctg ccacccgccc acccgccttg tgagatggaa	2113
ttgtaataaa ccacgccatg aggacaccgc cgcccgcctc ggcgcttcct ccaccgaaaa	2173
aaaaaaaaa aaaa	2187

FIG. 2

(SEQ ID NO: 2) Met Ser Ala Ile Gln Ala Ala Trp Pro Ser Gly Thr Glu Cys Ile Ala Lys Tyr Asn Phe His Gly Thr Ala Glu Gln Asp Leu Pro Phe Cys Lys Gly Asp Val Leu Thr Ile Val Ala Val Thr Lys Asp Pro Asn Trp Tyr Lys Ala Lys Asn Lys Val Gly Arg Glu Gly Ile Ile Pro Ala Asn Tyr Val Gln Lys Arg Glu Gly Val Lys Ala Gly Thr Lys Leu Ser Leu Met Pro Trp Phe His Gly Lys Ile Thr Arg Glu Gln Ala Glu Arg Leu Leu Tyr Pro Pro Glu Thr Gly Leu Phe Leu Val Arg Glu Ser Thr Asn Tyr Pro Gly Asp Tyr Thr Leu Cys Val Ser Cys Asp Gly Lys Val Glu His Tyr Arg Ile Met Tyr His Ala Ser Lys Leu Ser Ile Asp Glu Glu Val Tyr Phe Glu Asn Leu Met Gln Leu Val Glu His Tyr Thr Ser Asp Ala Asp Gly Leu Cys Thr Arg Leu Ile Lys Pro Lys Val Met Glu Gly Thr Val Ala Ala Gln Asp Glu Phe Tyr Arg Ser Gly Trp Ala Leu Asn Met Lys Glu Leu Lys Leu Leu Gln Thr Ile Gly Lys Gly Glu Phe Gly Asp Val Met Leu Gly Asp Tyr Arg Gly Asn Lys Val Ala Val Lys Cys Ile Lys Asn Asp Ala Thr Ala Gln Ala Phe Leu Ala Glu Ala Ser Val Met Thr Gln Leu Arg His Ser Asn Leu Val Gln Leu Leu Gly Val Ile Val Glu Glu Lys Gly Gly Leu Tyr Ile Val Thr Glu Tyr Met Ala Lys Gly Ser Leu Val Asp Tyr Leu Arg Ser Arg Gly Arg Ser Val Leu Gly Gly Asp Cys Leu Leu Lys Phe Ser Leu Asp Val Cys Glu Ala Met Glu Tyr Leu Glu Gly Asn Asn Phe Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Ser Glu Asp Asn Val Ala Lys Val Ser Asp Phe Gly Leu Thr Lys Glu Ala Ser Ser Thr Gln Asp Thr Gly Lys Leu Pro Val Lys Trp Thr Ala Pro Glu Ala Leu Arg Glu Lys Lys Phe Ser Thr Lys Ser Asp Val Trp Ser Phe Gly Ile Leu Leu Trp Glu Ile Tyr Ser Phe Gly Arg Val Pro Tyr Pro Arg Ile Pro Leu Lys Asp Val Val Pro Arg Val Glu Lys Gly Tyr Lys Met Asp Ala Pro Asp Gly Cys Pro Pro Ala Val Tyr Glu Val Met Lys Asn Cys Trp His Leu Asp Ala Ala Met Arg Pro Ser Phe Leu Gln Leu Arg Glu Gln Leu Glu His Ile Lys Thr His Glu Leu His Leu

FIG. 3

gcgg;	ID agcc	NO: aa g	3) gcac	acgg	g to	tgac	cctt	ggg	ccgg	ccc	ggag	caag	tg a	cacg	gaccg	60
gtcg	ccta	tc c	tgac	caca	g ca	aago	ggcc	cgg	agco	cgc	ggag	ıggga	icc t	gacg	ggggc	120
gtag	gcgc	cg g	aagg	ctgg	g gg	cccc	ggag	ccg	ggcc	ggc	gtgg	cccg	ag t	tccg	gtgag	180
cgga	cggc	gg c	gcgc	gcag	a tt	tgat	a at Me	g gg	rc tg .y Cy	c at	t aa le Ly	a ag	r Ly	a ga ⁄s Gl	a aac u Asn	234
aaa Lys 10	agt Ser	cca Pro	gcc Ala	att Ile	aaa Lys 15	tac Tyr	aga Arg	cct Pro	gaa Glu	aat Asn 20	act Thr	cca Pro	gag Glu	cct Pro	gtc Val 25	282
agt Ser	aca Thr	agt Ser	gtg Val	agc Ser 30	cat His	tat Tyr	gga Gly	gca Ala	gaa Glu 35	ccc Pro	act Thr	aca Thr	gtg Val	tca Ser 40	cca Pro	330
tgt Cys	ccg Pro	tca Ser	tct Ser 45	tca Ser	gca Ala	aag Lys	gga Gly	aca Thr 50	gca Ala	gtt Val	aat Asn	ttc Phe	agc Ser 55	agt Ser	ctt Leu	37.8
tcc Ser	atg Met	aca Thr 60	cca Pro	ttt Phe	gga Gly	gga Gly	tcc Ser 65	tca Ser	ggg Gly	gta Val	acg Thr	cct Pro 70	ttt Phe	gga Gly	ggt Gly	426
gca Ala	tct Ser 75	tcc Ser	tca Ser	ttt Phe	tca Ser	gtg Val 80	gtg Val	cca Pro	agt Ser	tca Ser	tat Tyr 85	cct Pro	gct Ala	ggt Gly	tta Leu	474
aca Thr 90	ggt Gly	ggt Gly	gtt Val	act Thr	ata Ile 95	ttt Phe	gtg Val	gcc Ala	tta Leu	tat Tyr 100	gat Asp	tat Tyr	gaa Glu	gct Ala	aga Arg 105	522
act Thr	aca Thr	gaa Glu	gac Asp	ctt Leu 110	tca Ser	ttt Phe	aag Lys	aag Lys	ggt Gly 115	gaa Glu	aga Arg	ttt Phe	caa Gln	ata Ile 120	att Ile	570
aac Asn	aat Asn	acg Thr	gaa Glu 125	gga Gly	gat Asp	tgg Trp	tgg Trp	gaa Glu 130	gca Ala	aga Arg	tca Ser	atc Ile	gct Ala 135	aca Thr	gga Gly	618
aag Lys	aát Asn	ggt Gly 140	tat Tyr	atc Ile	ccg Pro	agc Ser	aat Asn 145	tat Tyr	gta Val	gcg Ala	cct Pro	gca Ala 150	gat Asp	tcc Ser	att Ile	666
cag Gln	gca Ala 155	gaa Glu	gaa Glu	tgg Trp	tat Tyr	ttt Phe 160	Gly	aaa Lys	atg Met	GJĀ āāā	aga Arg 165	aaa Lys	gat Asp	gct Ala	gaa Glu	714
aga Arg 170	tta Leu	ctt Leu	ttg Leu	aat Asn	cct Pro 175	gga Gly	aat Asn	caa Gln	cga Arg	ggt Gly 180	Ile	ttc Phe	tta Leu	gta Val	aga Arg 185	762
gag Glu	agt Ser	gaa Glu	aca Thr	act Thr 190	Lys	ggt Gly	gct Ala	tat Tyr	tcc Ser 195	Leu	tct Ser	att Ile	cgt Arg	gat Asp 200	\mathtt{Trp}	810

FIG. 3 cont.

gat Asp	gag Glu	ata Ile	agg Arg 205	ggt Gly	gac Asp	aat Asn	gtg Val	aaa Lys 210	cac His	tac Tyr	aaa Lys	att Ile	agg Arg 215	aaa Lys	ctt Leu	858
gac Asp	aat Asn	ggt Gly 220	gga Gly	tac Tyr	tat Tyr	atc Ile	aca Thr 225	acc Thr	aga Arg	gca Ala	caa Gln	ttt Phe 230	gat Asp	act Thr	ctg Leu	906
cag Gln	aaa Lys 235	ttg Leu	gtg Val	aaa Lys	cac His	tac Tyr 240	aca Thr	gaa Glu	cat His	gct Ala	gat Asp 245	ggt Gly	tta Leu	tgc Cys	cac His	954
aag Lys 250	ttg Leu	aca Thr	act Thr	gtg Val	tgt Cys 255	cca Pro	act Thr	gtg Val	aaa Lys	cct Pro 260	cag Gln	act Thr	caa Gln	ggt Gly	cta Leu 265	1002
gca Ala	aaa Lys	gat Asp	gct Ala	tgg Trp 270	gaa Glu	atc Ile	cct Pro	cga Arg	gaa Glu 275	tct Ser	ttg Leu	cga Arg	cta Leu	gag Glu 280	gtt Val	1050
aaa Lys	cta Leu	gga Gly	caa Gln 285	gga Gly	tgt Cys	ttc Phe	ggc Gly	gaa Glu 290	gtg Val	tgg Trp	atg Met	gga Gly	aca Thr 295	tgg Trp	aat Asn	1098
gga Gly	acc Thr	acg Thr 300	aaa Lys	gta Val	gca Ala	atc Ile	aaa Lys 305	aca Thr	cta Leu	aaa Lys	cca Pro	ggt Gly 310	aca Thr	atg Met	atg Met	1146
cca Pro	gaa Glu 315	gct Ala	tt <i>c</i> Phe	ctt Leu	caa Gln	gaa Glu 320	gct Ala	cag Gln	ata Ile	atg Met	aaa Lys 325	aaa Lys	tta Leu	aga Arg	cat His	1194
gat Asp 330	aaa Lys	ctt Leu	gtt Val	cca Pro	cta Leu 335	tat Tyr	gct Ala	gtt Val	gtt Val	tct Ser 340	gaa Glu	gaa Glu	cca Pro	att Ile	tac Tyr 345	1242
att Ile	gtc Val	act Thr	gaa Glu	ttt Phe 350	atg Met	tca Ser	aaa Lys	.gga Gly	agc Ser 355	tta Leu	tta Leu	gat Asp	ttc Phe	ctt Leu 360	aag Lys	1290
gaa Glu	gga Gly	gat Asp	gga Gly 365	aag Lys	tat Tyr	ttg Leu	aag Lys	ctt Leu 370	cca Pro	cag Gln	ctg Leu	gtt Val	gat Asp 375	atg Met	gct Ala	1338
gct Ala	cag Gln	att Ile 380	Ala	gat Asp	ggt Gly	atg Met	gca Ala 385	tat Tyr	att Ile	gaa Glu	aga Arg	atg Met 390	aac Asn	tat Tyr	att Ile	1386
cac His	cga Arg 395	Asp	ctt Leu	cgg Arg	gct Ala	gct Ala 400	Asn	att Ile	ctt Leu	gta Val	gga Gly 405	gaa Glu	aat Asn	ctt Leu	gtg Val	1434
tgc Cys 410	Lys	ata Ile	gca Ala	gac Asp	ttt Phe 415	Gly	tta Leu	gca Ala	agg Arg	tta Leu 420	Ile	gaa Glu	gac Asp	aat Asn	gaa Glu 425	1482
tac Tyr	aca Thr	gca Ala	aga Arg	caa Gln 430	ggt	gca Ala	aaa Lys	ttt Phe	cca Pro 435	Ile	aaa Lys	tgg Trp	aca Thr	gct Ala 440	Pro	1530

FIG. 3 cont.

Phe Gly Ile Leu Gln Thr Glu Leu Val Thr Lys Gly Arg Val Pro Tyr 460 cca ggt atg gtg aac cgt gaa gta cta gaa caa gtg gag cga gga tac Pro Gly Met Val Asn Arg Glu Val Leu Glu Gln Val Glu Arg Gly Tyr 475 agg atg ccg tgc cct cag ggc tgt cca gaa tcc ctc cat gaa ttg atg Arg Met Pro Cys Pro Gln Gly Cys Pro Glu Ser Leu His Glu Leu Met 500 aat ctg tgt tgg aag aag gac cct gat gaa aga cca aca ttt gaa tat 1770 Asn Leu Cys Trp Lys Lys Asp Pro Asp Glu Arg Pro Thr Phe Glu Tyr 510 att cag tcc ttc ttg gaa gac tac ttc act gct aca gag cca cag tac I818 Ile Gln Ser Phe Leu Glu Asp Tyr Phe Thr Ala Thr Glu Pro Gln Tyr 535	gaa Glu	gct Ala	gca Ala	ctg Leu 445	tat Tyr	ggt Gly	cgg Arg	ttt Phe	aca Thr 450	ata Ile	aag Lys	tct Ser	gat Asp	gtc Val 455	tgg Trp	tca Ser	1578
Pro Gly Met Val Asn Arg Glu Val Leu Glu Glu Val Glu Arg Gly Tyr 485 agg atg ccg tgc cct cag ggc tgt cca gaa tcc ctc cat gaa ttg atg Arg Met Pro Cys Pro Gln Gly Cys Pro Glu Ser Leu His Glu Leu Met 505 aat ctg tgt tgg aaag aag gac cct gat gaa aag cca aca ttt gaa tat Asn Leu Cys Trp Lys Lys Asp Pro Asp Glu Arg Pro Thr Phe Glu Tyr 510 att cag tcc ttc ttg gaa gac tac ttc act gct aca gag cca cag tac lle Gln Ser Phe Leu Glu Asp Tyr Phe Thr Ala Thr Glu Pro Gln Tyr 520 att cag cca gga gaa aat tta taattcaagt agcctattt atatgcacaa 1866 Gln Pro Gly Glu Asn Leu 540 atctgccaaa atataaagaa cttgtgtaga ttttctacag gaatcaaaag aagaaaatct 1920 cttttactct gcatgtttt aaaggtaaac tggaatcca gaattggttg cacaaaacca 1980 ctttttttc cccaagtatt aaactctaat gtaccaatga tggattatc agcgtatttc 2040 agggtccaaa caaaatagag ctaagatact gatgacagtg tgggtgacag catggtaatg 210 aaggacagtg aggctoctge ttattataa atcattcct ttctttttt cccaaagtc 226 agaattgctc aaagaaaatt atttattgtt acagataaaa cttgagaga aaaaacta 222 accataataa aactcaaaat taaggaatat catggacca aataatcca ttccagtttt 228 ttaaagttc ttgcatttat tattccaaa agtttttct aagttaaaca gtcagtate 234 aaccataataa tatgcttct tttgcatgga catggaccag gttttcaaa aggaatataa 240 acaggacctc aaacttgatt catggaactg aagaataga gaattaacc 240 agagtactaata tattcatgtt catggaactg aagaataga aacttttca cttcagtcc 252 tttctgaaga gtttgactta gaataatga ggaaataa aattttca catggaact aactttaa catttcat ttctgaatga catggaga 264 agtacattaa tattcatgtt catggaactg aagaataag aacttttca cttcagtcc 252 tttctgaaga gtttgactta gaataatga ggtaactaga aagtaggta accttgatag 258 aggttgcatt gatttttaa ggcaatata aattgaacca attttaaaag ccctttttta 270 aaaactagac ttcgtactg gagtattgct tatatgtcc tatatggaac ggtgccaca 276 aagactagac ttcgtactg gagtattgct tatatgtcc tatatgggat ggtgccaca 276 aagactagac ttcgtactg gagtattgct tatatgtcc tatatggaac ggtgccaca 276 aagactaga gtttgactt gagtattgct tatatgtcc tatatggca 276 aagactaga gtttgactt gagtattgct tatatgtcc tatatgggat ggtgccaca 276	ttt Phe	gga Gly	Ile	ctg Leu	caa Gln	aca Thr	gaa Glu	Leu	gta Val	aca Thr	aag Lys	Gly	Arg	gtg Val	cca Pro	tat Tyr	1626
Arg Met Pro Cys Pro Gln Gly Cys Pro Glu Ser Leu His Glu Leu Met 490 aat ctg tgt tgg aag aag gac cct gat gaa aag acc a cca act ttt gaat 1770 Asn Leu Cys Trp Lys Lys Asp Pro Asp Glu Arg Pro Thr Phe Glu Tyr 510 att cag tcc ttc ttg gaa gac tac ttc act gct aca gag cca cag tac 11e Gln Ser Phe Leu Glu Asp Tyr Phe Thr Ala Thr Glu Pro Gln Tyr 525 cag cca gga gaa aat tta taattcaagt agcctatttt atatgcacaa 1860 Acttgccaaa atataaagaa cttgtgtaga ttttctacag gaatcaaaag aagaaaatct 1920 Acttgccaaa atataaagaa cttgtgtaga ttttctacag gaatcaaaag aagaaaatc 1920 Actttttttc cccaagtatt aaactctaat gtaccaatga tgagttg cacaaaacca 1980 Agggtccaaa caaaatagag ctaagatact gatgacagtg tgggtgacag catggtattc 2040 Aagagtgagtg aggctcctge ttattataa atcattcct ttctttttt ccccaaagtc 216 Agaattgctc aaagaaaatt atttattgtt acagataaaa cttgagaga aaaaactat 222 Accataataa aatctaaaat taatggaatat catgggaca aataattcca ttccagtttt 228 Aaacttaata tatgctttct tttgcatgga catgggccag gttttcaaa aggaatata 240 Acaggacctc aaacttgatt aaatgttaga ccacagaagt ggaatttgaa agtataatc 246 Agatctaata tattcatgtt catggaactg aaagaataag aacttttca cttcagtcc 252 Accataataa tattcatgtt catggaactg aaagaataag aacttttca cttcagtcc 254 Aatcttaata tattcatgtt catggaactg aaagaataag aacttttca cttcagtcc 254 Aggttgcatt gatttttaa ggcaatatat aattgaact actggacca caaagaggaaccc 264 Aggttgcatt gatttttaa ggcaatatat aattgaacc accagaagt aagtgagtta acttgatag 258 Aggttgcatt gatttttaa ggcaatatat aattgaacc accagaagt aagtgagtta acttgtatg 258 Aggttgcatt gatttttaa ggcaatatat aattgaacc actggacca attttaaaag ccctttttta 270 Aaaaccagac ttcgtactgt gagtattgct tatatgccc aaaccaga 276 Aggttgcatt gatttttaa ggcaatatat aattgaacc accagaagt accacagaagt accacagaagt accacagaagt accacagaaccacacacacacacacacacacacacacac	cca Pro	Ğly	atg Met	gtg Val	aac Asn	cgt Arg	Glu	gta Val	cta Leu	gaa Glu	caa Gln	Val	gag Glu	cga Arg	gga Gly	tac Tyr	1674
Asn Leu Cys Trp Lys Lys Asp Pro Asp Glu Arg Pro Thr Phe Glu Tyr 510 att cag tcc tr ttg gaa gac tac trace act gct aca gag cca cag tac 1818 [Ile Gln Ser Phe Leu Glu Asp Tyr Phe Thr Ala Thr Glu Pro Gln Tyr 525 cag cca gga gaa aat tta taattcaagt agcctattt atatgcacaa 1866 [In Pro Gly Glu Asn Leu 540] atctgccaaa atataaagaa cttgtgtaga ttttctacag gaatcaaaag aagaaaatct 1926 [Ite ttttctct gcatgtttt aatggtaaac tggaatcca gatatggtg cacaaaacca 1986 [Ite ttttttc cccaagtatt aaactctaat gtaccaatga tggattatc agcgtattc 2046] agggtccaaa caaaatagag ctaagatact gatgacagtg tgggtgacag catggtaatg 210 aaggacagtg aggctcctgc ttatttataa atcatttcct ttctttttt ccccaaagtc 216 agaattgctc aaagaaaatt attattgtt acaggaacaa catggagat aaaaagctat 222 accataataa aatctaaaat taatggaatat catgggacca aataattcca ttccagttt 228 ttaaagttc ttgcatttat tattctcaaa agtttttct aagtaaaca gtcagtatgc 234 aatcttaata tatgctttct tttgcatgga catgggcag gttttcaaa aggaatataa 240 acaggatctc aaacttgat aaatgtaga ccacagaagt ggaatttgaa agtataatg 240 acaggatctc aaacttgat catggaactg aaagaataag aacttttca cttcagtcct 252 tttctgaaga gtttgactta gaataatgaa ggtaactaga aagtgagtta atcttgatg 258 aggttgcatt gatttttaa ggcaatatat aattgaaact actgtccaat caaaggggaa 264 atgttttgat ctttagatag catgcaaagt aagaccage attttaaaag ccctttttta 270 aaaactagac ttcgtactg gagtattgct tatatgtcct tatagggatg ggtgccacaa 276	Arg	atg Met	ccg Pro	tgc Cys	cct Pro	Gln	ggc Gly	tgt Cys	cca Pro	gaa Glu	Ser	ctc Leu	cat His	gaa Glu	ttg Leu	Met	1722
cag cca gga gaa aat tta taattcaagt agcctattt atatgcacaa 1866 Clin Pro Cly Cly Cly Asn Leu atctgccaaa atataaagaa cttgtgtaga ttttctacag gaatcaaaag aagaaaatct 1920 tctttactct gcatgtttt aatggtaaac tggaatcca gatatggttg cacaaaacca 1986 ctttttttc cccaagtat aaactctaat gtaccaatga tgaatttatc agcgtatttc 2046 agggtccaaa caaaatagag ctaagatact gatgacagtg tgggtgacag catggtaatg 210 aaggacagtg aggctcctge ttatttataa atcatttcct ttcttttt ccccaaagtc 216 agaattgctc aaagaaaatt atttattgtt acagataaaa cttgagagat aaaaagctat 222 accataataa aatctaaaat taatgaatat catgggacca aataattcca ttccagtttt 228 ttaaagtttc ttgcatttat tattctcaaa agtttttct aagttaaaca gtcagtatgc 234 aatcttaata tatgctttct tttgcatgga catgggccag gttttcaaa aggaatataa 240 acaggatctc aaacttgatt aaatgttaga ccacagaagt ggaatttgaa agtataatgc 246 agtacattaa tattcatgtt catggaactg aaagaataag aacttttca cttcagtcct 252 tttctgaaga gtttgactta gaataatgaa ggtaactaga aagtgagtta atcttgatg 258 aggttgcatt gatttttaa ggcaatata aattgaaact actgtccaat caaaggggaa 264 atgttttgat ctttagatag catgcaaagt aagaccagc attttaaaag ccctttttta 270 aaaaactagac ttcgtactgt gagtattgct tatatgcct tatggggatg ggtgccacaa 276	aat Asn	ctg Leu	tgt Cys	tgg Trp	Lys	aag Lys	gac Asp	cct Pro	gat Asp	Glu	aga Arg	cca Pro	aca Thr	ttt Phe	Glu	tat Tyr	1770
atctgccaaa atataaagaa cttgtgtaga ttttctacag gaatcaaaag aagaaaatct 1926 tctttactct gcatgtttt aatggtaaac tggaatccca gatatggttg cacaaaacca 1986 ctttttttc cccaagtatt aaactctaat gtaccaatga tggattatc agcgtattc 2046 agggtccaaa caaaatagag ctaagatact gatgacagtg tgggtgacag catggtaatg 210 aaggacagtg aggctcctge ttatttataa atcatttcct ttctttttt ccccaaagtc 216 agaattgctc aaagaaaatt atttattgtt acagataaaa cttgagagat aaaaagctat 222 accataataa aatctaaaat taatggaatat catgggacca aataattcca ttccagtttt 228 ttaaagtttc ttgcatttat tattctcaaa agtttttct aagttaaaca gtcagtatgc 234 aatcttaata tatgcttct tttgcatgga catgggccag gttttcaaa aggaatataa 240 acaggatctc aaacttgatt aaatgtaga ccacagaagt ggaatttgaa agtataatgc 246 agtacattaa tattcatgtt catggaactg aaagaataag aacttttca cttcagtcct 252 tttctgaaga gtttgactta gaataatgaa ggtaactaga aagtgagtta atcttgtatg 258 aggttgcatt gatttttaa ggcaatatat aattgaaact actgtccaat caaaggggaa aaaactagac ttctagatag catgcaaagt aagacccage attttaaaag ccctttttta 270 aaaactagac ttcgtactgt gagtattgct tatatgtcct tatatgtcct tatagggatg ggtgccacaa 276	att Ile	cag Gln	tcc Ser	Phe	ttg Leu	gaa Glu	gac Asp	tac Tyr	Phe	act Thr	gct Ala	aca Thr	gag Glu	Pro	Gln	tac Tyr	1818
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aaaactagac ttcgtactgt gagtattgct tatatgtcct tatgggggatg ggtgccacaa 276																	
	ata	agaaa	aata	tgad	ccaga	atc a	aggga	actto	ga at	gcad	tttt	gct	cato	ggtg	aata	tagatg	2826

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FIG. 3 cont.

aacagagagg aaaatgtatt taaaagaaat acgagaaaag aaaatgtgaa agttttacaa 2886 gttagaggga tggaaggtaa tgtttaatgt tgatgtcatg gagtgacaga atggctttgc 2946 tggcactcag agetectcae ttagetatat tetgagaett tgaagagtta taaagtataa 3006 ctataaaact aatttttctt acacactaaa tgggtatttg ttcaaaataa tgaagttatg 3066 gcttcacatt cattgcagtg ggatatggtt tttatgtaaa acatttttag aactccagtt 3126 ttcaaatcat gtttgaatct acattcactt ttttttgttt tcttttttga gacggagtct 3186 cgctctgccg cccaggctgg agtgcagtgg cgcgatctcg gctcactgca agctctgcct 3246 cccaggttca caccattctc ctgcctcagc ctcccgagta gctgggacta caggtgccca 3306 ccaccacgcc tggctagttt tttgtatttt tagtagagac gcagtttcac cgtgttagcc 3366 aggatggtct cgatctcctg accttgtgat ctgcccgcct cggcctccca aagtgctggg 3426 attacaggtg tgagccaccg cgcccagcct acattcactt ctaaagtcta tgtaatggtg 3486 gtcatttttt cccttttaga atacattaaa tggttgattt ggggaggaaa acttattctg 3546 aatattaacg gtggtgaaaa ggggacagtt tttaccctaa agtgcaaaag tgaaacatac 3606 aaaataagac taatttttaa gagtaactca gtaatttcaa aatacagatt tgaatagcag 3666 cattagtggt ttgagtgtct agcaaaggaa aaattgatga ataaaatgaa ggtctggtgt 3726 atatgtttta aaatactctc atatagtcac actttaaatt aagccttata ttaggcccct 3786 ctattttcag gatataattc ttaactatca ttatttacct gattttaatc atcagattcg 3846 aaattctgtg ccatggcgta tatgttcaaa ttcaaaccat ttttaaaatg tgaagatgga 3906 cttcatgcaa gttggcagtg gttctggtac taaaaattgt ggttgttttt tctgtttacg 3966 taacctgctt agtattgaca ctctctacca agagggtctt cctaagaaga gtgctgtcat 4026 tatttcctct tatcaacaac ttgtgacatg agatttttta agggctttat gtgaactatg 4086 atattgtaat ttttctaagc atattcaaaa gggtgacaaa attacgttta tgtactaaat 4146 ctaatcagga aagtaaggca ggaaaagttg atggtattca ttaggtttta actgaatgga 4206 gcagttcctt atataataac aattgtatag tagggataaa acactaacaa tgtgtattca 4266 ttttaaattg ttctgtattt ttaaattgcc aagaaaaaca actttgtaaa tttggagata 4326 ttttccaaca gcttttcgtc ttcagtgtct taatgtggaa gttaaccctt accaaaaaag 4386 gaagttggca aaaacagcct tctagcacac ttttttaaat gaataatggt agcctaaact 4446 taatattttt ataaagtatt gtaatattgt tttgtggata attgaaataa aaagttctca 4506 4517 ttgaatgcac c

FIG. 4

(SEQ ID NO: 4)

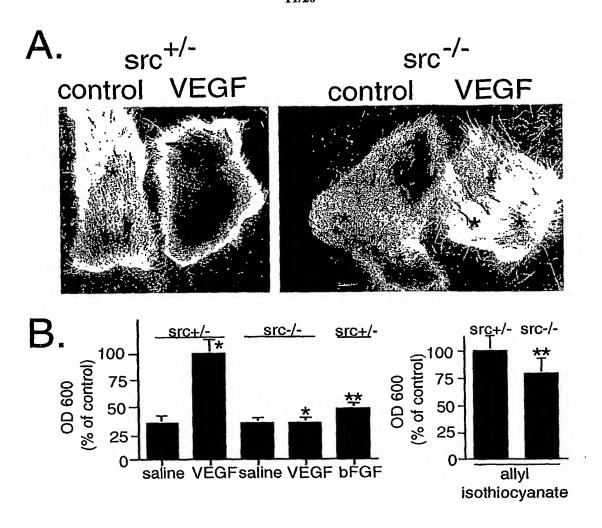
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	Pro	Glu	Asn 20	Thr	Pro	Glu	Pro	Val 25	Ser	Thr	Ser	Val	Ser 30	His	Tyr
Gly	Ala	Glu 35		Thr	Thr	Val	Ser 40	Pro	Cys	Pro	Ser	Ser 45	Ser	Ala	Lys
Gly	Thr 50		Val	Asn	Phe	Ser 55	Ser	Leu	Ser	Met	Thr 60	Pro	Phe	Gly	Gly
Ser 65	Ser	Gly	Val	Thr	Pro 70		Gly	Gly	Ala	Ser 75	Ser	Ser	Phe	Ser	Val 80
Val	Pro	Ser	Ser	Tyr 85		Ala	Gly	Leu	Thr 90		Gly	Val	Thr	Ile 95	Phe
Val	Ala	Leu	Tyr 100		Tyr	Glu	Ala	Arg 105	Thr	Thr	Glu	Asp	Leu 110	Ser	Phe
Lys	Lys	Gly 115	Glu	Arg	Phe	Gln	Ile 120		Asn	Asn	Thr	Glu 125	Gly	Asp	Trp
Trp	Glu 130	Ala	Arg	Ser	Ile	Ala 135	Thr	Gly	Lys	Asn	Gly 140	Tyr	Ile	Pro	Ser
145	Tyr				150					155				Tyr	160
Gly	Lys	Met	Gly	Arg 165	Lys	Asp	Ala	Glu	Arg 170	Leu	Leu	Leu	Asn	Pro 175	Gly
Asn	Gln	Arg	Gly 180	Ile	Phe	Leu	Val	Arg 185	Glu	Ser	Glu	Thr	Thr 190	Lys	Gly
Ala	Tyr	Ser 195		Ser	Ile	Arg	Asp 200	Trp	Asp	Glu	Ile	Arg 205	Gly	Asp	Asn
Val	Lуs 210	His	Tyr	Lys	Ile	Arg 215	Lys	Leu	Asp	Asn	Gly 220	Gly	Tyr	Tyr	Ile
Thr 225	Thr	Arg	Ala	Gln	Phe 230	Asp	Thr	Leu	Gln	Lys 235	Leu	Val	Lys	His	Tyr 240
Thr				245					250					Cys 255	
		_	260					265					270	Glu	
		275					280					285		Cys	
_	290		_			295					300			Ala	
305					310					315				Gln	320
				325					330					Leu 335	
			340					345					350		
		355					360					365		Tyr	
_	370					375					380			Gly	
385					390					395				Ala	400
Asn	Ile			405					410					Phe 415	
Leu	. Ala	Arg	Leu 420		Glu	Asp	Asn	Glu 425		Thr	Ala	Arg	Gln 430	Gly	Ala

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FIG. 4 cont.

Lys	Phe	Pro 435	Ile	Lys	Trp	Thr	Ala 440	Pro	Glu	Ala	Ala	Leu 445	Tyr	Gly	Arg
Phe	Thr 450	Ile	Lys	Ser	Asp	Val 455	Trp	Ser	Phe	Gly	Ile 460	Leu	Gln	Thr	Glu
Leu 465	Val	Thr	Lys	Gly	Arg 470	Val	Pro	Tyr	Pro	Gly 475	Met	Val	Asn	Arg	Glu 480
Val	Leu	Glu	Gln	Val 485	Glu	Arg	Gly	Tyr	Arg 490	Met	Pro	Cys	Pro	Gln 495	Gly
Cys	Pro	Glu	Ser 500	Leu	His	Glu	Leu	Met 505	Asn	Leu	Cys	Trp	Lys 510	Lys	Asp
Pro	Asp	Glu 515	Arg	Pro	Thr	Phe	Glu 520	Tyr	Ile	Gln	Ser	Phe 525	Leu	Glu	Asp
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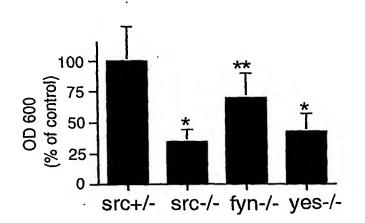


FIG. 5

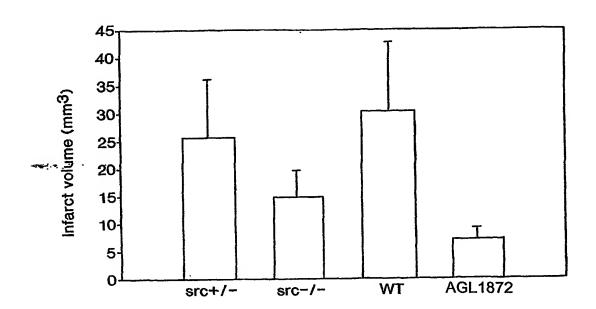
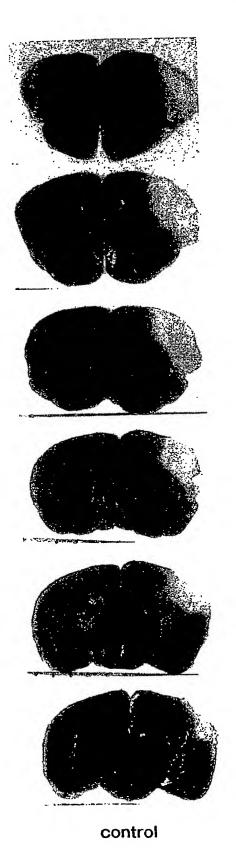


FIG. 6

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FIG. 7

FIG. 8

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OCONH₂

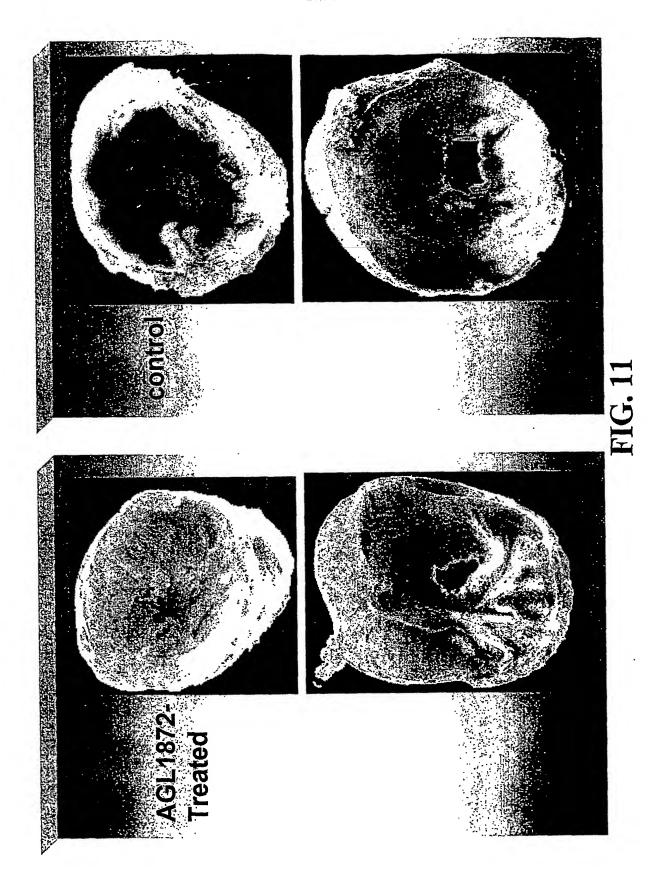
FIG. 9

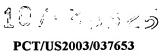
ĊH₃

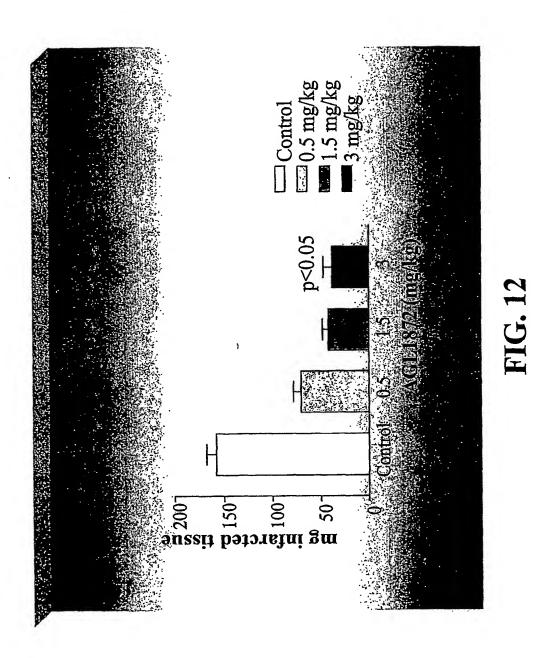
PD173955

FIG. 10









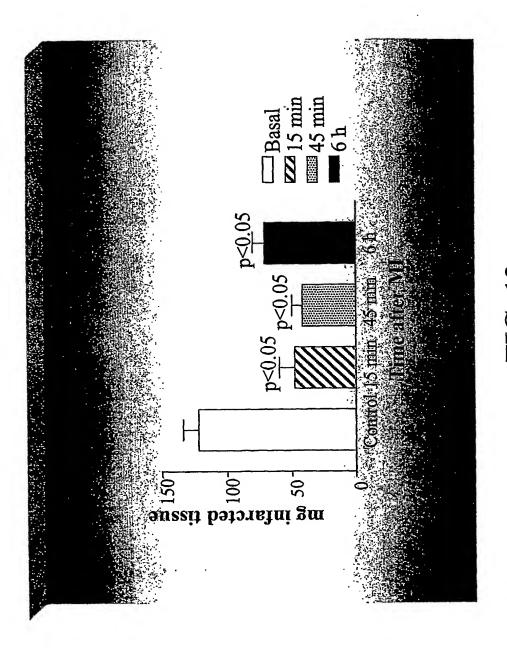
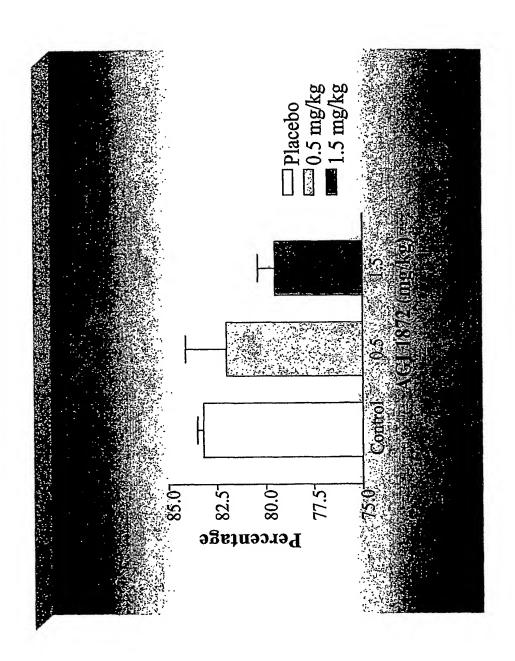


FIG. 13





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